

EBI Dbfetch

ID AA689524 standard; mRNA; EST; 544 BP.
 XX
 AC AA689524;
 XX
 SV AA689524.1
 XX
 DT 19-DEC-1997 (Rel. 54, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 3)
 XX
 DE ns66e01.r1 NCI_CGAP_Pr22 Homo sapiens cDNA clone IMAGE:1188600 5' similar
 DE to SW:CNI_DROME P49858 CORNICHON PROTEIN. ;, mRNA sequence.
 XX
 KW EST.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-544
 RA NCI-CGAP;
 RT "National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor
 RT Gene Index <http://www.ncbi.nlm.nih.gov/ncicgap>;
 RL Unpublished.
 XX
 DR RZPD; [IMAGp998K012985](#).
 DR UNILIB; [34](#); 910.
 XX
 CC Insert Length: 484 Std Error: 0.00
 CC Seq primer: -28m13 rev1 ET from Amersham
 CC High quality sequence stop: 499.
 XX
 CC On Nov 6, 1997 this sequence version replaced gi:947554.
 CC Contact: Robert Strausberg, Ph.D.
 CC Tel: (301) 496-1550
 CC Email: Robert_Strausberg@nih.gov
 CC Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.
 CC Emmert-Buck, M.D., Ph.D.
 CC cDNA Library Preparation: M. Bento Soares, Ph.D.
 CC cDNA Library Arrayed by: Greg Lennon, Ph.D.
 CC DNA Sequencing by: Washington University Genome Sequencing Center
 CC Clone distribution: NCI-CGAP clone distribution information can be
 CC found through the I.M.A.G.E. Consortium/LLNL at:
 CC www-bio.llnl.gov/bbrp/image/image.html
 XX
 FH Key Location/Qualifiers
 FH
 FT source 1..544
 FT /db_xref="taxon:9606"
 FT /db_xref="RZPD:IMAGp998K012985"
 FT /db_xref="UNILIB:34"

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FT      /mol_type="mRNA"
FT      /note="Organ: prostate; Vector: pT7T3D-Pac (Pharmacia) with
FT      a modified polylinker; 1st strand cDNA was prepared from
FT      normal prostate bulk tissue, and was then primed with a Not
FT      I - oligo(dT) primer. Double-stranded cDNA was ligated to
FT      Eco RI adaptors (Pharmacia), digested with Not I and cloned
FT      into the Not I and Eco RI sites of the modified pT7T3
FT      vector. Library is normalized, and was constructed by
FT      Bento Soares and M. Fatima Bonaldo. "
FT      /sex="male"
FT      /organism="Homo sapiens"
FT      /clone="IMAGE:1188600"
FT      /clone_lib="NCI_CGAP_Pr22"
FT      /tissue_type="normal prostate"
FT      /lab_host="DH10B"
XX

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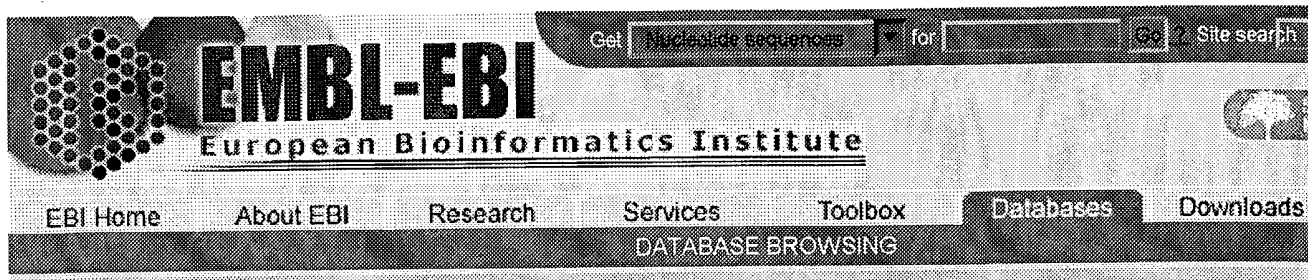
SQ      Sequence 544 BP; 165 A; 91 C; 103 G; 185 T; 0 other;
      gtgatgagtg gccaggactc tatgacccta caaccatcat gaatgcagat attctagcat      60
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      tatatggcat gatctatggt ttggtgagct cttagaacaa cacacagaag aattgggtcca      180
      gttaagtgca tgcaaaaagc caccaaatga agggattcta tccagcaaga tctgtgccaa      240
      gagtagcctg tggaatctga tcagttactt taaaaaatga ctcttatctt tttaaatggt      300
      tccacatttt tgcttggtga aagactgttt tcatatgtta tactcagata aagattttta      360
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      gcacttcotta aggaacagcc ataatectct gaatgatgca ttaattactg actgtcctag      480
      tacattggaa gcttttgttt ataggaactt gtagggctca ttttggtttc attgaaacag      540
      tatc

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EBI Dbfetch

ID AA902726 standard; mRNA; EST; 500 BP.
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 AC AA902726;
 XX
 SV AA902726.1
 XX
 DT 09-APR-1998 (Rel. 55, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 3)
 XX
 DE ok71h07.s1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1519453 3', mRNA
 DE sequence.
 XX
 KW EST.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-500
 RA NCI-CGAP;
 RT "National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor
 RT Gene Index <http://www.ncbi.nlm.nih.gov/ncicgap>;
 RL Unpublished.
 XX
 DR RZPD; [IMAGp998D143847](#).
 DR UNILIB; [46](#); 1050.
 XX
 CC Insert Length: 1413 Std Error: 0.00
 CC Seq primer: -40m13 fwd. ET from Amersham
 CC High quality sequence stop: 375.
 XX
 CC On Jan 9, 1998 this sequence version replaced gi:936894.
 CC Contact: Robert Strausberg, Ph.D.
 CC Tel: (301) 496-1550
 CC Email: Robert.Strausberg@nih.gov
 CC Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
 CC Emmert-Buck, M.D., Ph.D.
 CC cDNA Library Preparation: M. Bento Soares, Ph.D.
 CC cDNA Library Arrayed by: Greg Lennon, Ph.D.
 CC DNA Sequencing by: Washington University Genome Sequencing Center
 CC Clone distribution: NCI-CGAP clone distribution information can be
 CC found through the I.M.A.G.E. Consortium/LLNL at:
 CC www-bio.llnl.gov/bbrp/image/image.html
 XX
 FH Key Location/Qualifiers
 FH
 FT source 1..500
 FT /db_xref="taxon:9606"
 FT /db_xref="RZPD:IMAGp998D143847"
 FT /db_xref="UNILIB:46"

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FT      /mol_type="mRNA"
FT      /note="Vector: pT7T3D-Pac (Pharmacia) with a modified
FT      polylinker; 1st strand cDNA was prepared from 3 pooled germ
FT      cell tumors, and was then primed with a Not I - oligo(dT)
FT      primer. Double-stranded cDNA was ligated to Eco RI adaptors
FT      (Pharmacia), digested with Not I and cloned into the Not I
FT      and Eco RI sites of the modified pT7T3 vector. Library is
FT      normalized. Library was constructed by Bento Soares and M.
FT      Fatima Bonaldo. "
FT      /organism="Homo sapiens"
FT      /clone="IMAGE:1519453"
FT      /clone_lib="NCI_CGAP_GC4"
FT      /tissue_type="pooled germ cell tumors"
FT      /lab_host="DH10B"

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XX
SQ      Sequence 500 BP; 180 A; 81 C; 66 G; 173 T; 0 other;
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      ctcatcagcag ttgaaaaatc tgtagtacct gcaaattcctt aagaatacct ttaccaccag      120
      attagaacag taagcataat aaccaatttc ttaataagta atgtcttaca aataaaaaaca      180
      catttaaaat agctttaaat gcattcttca caagtaattc agcatatatt tttatatcat      240
      gtttacttat gcttaagaat taaagcaagt atatttatta ctctgatgga aatatgggaa      300
      atctctcatt catgcaatat acagggataa tattcaagcg aagggaaaat tcccgcgttt      360
      tatttttgta aatgtatcca tatataatca tcgacatgac agatgaggaa acccatgaag      420
      tttcccacta gtcagatata cattttcact tcatcagaag cacctgatat ctacagctaa      480
      ttataatta gatactgttt

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